Sequence Listing

<110> Chen, Jian Filvaroff, Ellemane Fong, Sherman Goddard, Audrey Godowski, Paul L. Grimaldi, J.Christopher Gurney, Austin Li, Hanzhong Hillan, Kenneth J. Hymowitz, Sarah Tumas, Daniel Starovasnik, Melissa. VanLookeren, Menno Vandlen, Richard Watanabe, Colin Williams, P.Mickey Wood, William Yansura, Daniel

- <120> IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
- <130> P1381R1C1P4(US)
- <140> US 10/000,157
- <141> 2001-10-30
- <150> 60/085579
- <151> 1998-05-15
- <150> 60/113621
- <151> 1998-12-23
- <150> 60/130232
- <<del>151> 1999-04-21</del>
- <150> 60/131022
- <151> 1999-04-26
- <150> 60/134287
- <151> 1999-05-14
- <150> 60/138387
- <151> 1999-06-09
- <150> 60/172096
- <151> 1999-12-23
- <150> 60/175481
- <151> 2000-01-11
- <150> 60/191007
- <151> 2000-03-21
- <150> 60/213807

- <151> 2000-06-22
- <150> 60/242837
- <151> 2000-10-24
- <150> 60/244072
- <151> 2000-10-26
- <150> 60/253646
- <151> 2000-11-28
- <150> 09/311832
- <151> 1999-05-14
- <150> 09/380138
- <151> 1999-08-25
- <150> 09/380142
- <151> 1999-08-25
- <150> 09/644848
- <151> 2000-08-22
- <150> 09/747259
- <151> 2000-12-20
- <150> 09/816744
- <151> 2001-03-22
- <150> 09/854208
- <151> 2001-05-10
- <150> 09/854280
- <151> 2001-05-10
- <150> 09/874503
- <151> 2001-06-05
- <150> 09/908827
- <151> 2001-07-18
- <150> 09/918585
- <151> 2001-07-30
- <150> 09/929404
- <151> 2001-08-13
- <150> 09/931836
- <151> 2001-08-16
- <150> PCT/US99/05028
- <151> 1999-03-08

	1	
<150>	PCT/US99/10733	3

<151> 1999-05-14

<150> PCT/US99/31274

<151> 1999-12-30

<150> PCT/US00/04341

<151> 2000-02-18

<150> PCT/US00/05601

<151> 2001-03-01

<150> PCT/US00/05841

<151> 2000-03-02

<150> PCT/US00/07532

<151> 2000-03-21

<150> PCT/US00/15264

<151> 2000-06-02

<150> PCT/US00/23328

<151> 2000-08-24

<150> PCT/US00/30873

<151> 2000-11-10

<150> PCT/US00/32678

<151> 2000-12-01

<150> PCT/US00/34956

<151> 2000-12-20

<150> PCT/US01/06520

<151> 2001-02-28

<150> PCT/US01/17800

<151> 2001-06-01

<150> PCT/US01/19692

<151> 2001-06-20

<150> PCT/US01/21066

<151> 2001-06-29

<150> PCT/US01/21735

<151> 2001-07-09

<160> 39

<210> 1

 $\frac{\langle 211 \rangle}{\langle 212 \rangle} \frac{687}{DNA}$ 

## <213> Homo Sapien

education of the state of the s

<210> 2 <211> 180 <212> PRT <213> Homo Sapien

Met Asp Trp Pro His Asn Leu Leu Phe Leu Leu Thr Ile Ser Ile 15

Phe Leu Gly Leu Gly Gln Pro Arg Ser Pro Lys Ser Lys Arg Lys 30

Gly Gln Gly Arg Pro Gly Pro Leu Ala Pro Gly Pro His Gln Val 45

Pro Leu Asp Leu Val Ser Arg Met Lys Pro Tyr Ala Arg Met Glu 60

Glu Tyr Glu Arg Asn Ile Glu Glu Met Val Ala Gln Leu Arg Asn 75

Ser Ser Glu Leu Ala Gln Arg Lys Cys Glu Val Asn Leu Gln Leu 80

Trp Met Ser Asn Lys Arg Ser Leu Ser Pro Trp Gly Tyr Ser Ile 105

	· ·	
•	Asn His Asp Pro Ser Arg Ile Pro Val Asp Leu Pro Glu Ala Arg 110 115 120	
	Cys Leu Cys Leu Gly Cys Val Asn Pro Phe Thr Met Gln Glu Asp 125 130 135	
	Arg Ser Met Val Ser Val Pro Val Phe Ser Gln Val Pro Val Arg 140 145 150	
	Arg Arg Leu Cys Pro Pro Pro Pro Arg Thr Gly Pro Cys Arg Gln 155 160 165	
	Arg Ala Val Met Glu Thr Ile Ala Val Gly Cys Thr Cys Ile Phe 170 175 180	
•	<pre></pre>	
	<400> 3 gccaggtgtg caggccgctc caagcccagc ctgccccgct gccgccacca 50	
	tgacgeteet ecceggeete etgtttetga eetggetgea cacatgeetg 100	
	gcccaccatg acccctccct cagggggcac ccccacagtc acggtacccc 150	
	acactgctac tcggctgagg aactgcccct cggccaggcc ccccacacc 200	
	tgctggctcg aggtgccaag tgggggcagg ctttgcctgt agccctggtg 250	
	tccagcctgg aggcagcaag ccacaggggg aggcacgaga ggccctcagc 300	
	tacgacccag tgcccggtgc tgcggccgga ggaggtgttg gaggcagaca 350	
	cccaccagcg ctccatctca ccctggagat accgtgtgga cacggatgag 400	
	gaccgctatc cacagaagct ggccttcgcc gagtgcctgt gcagaggctg 450	
	tatcgatgca cggacgggcc gcgagacagc tgcgctcaac tccgtgcggc 500	
	tgctccagag cctgctggtg ctgcgccgcc ggccctgctc ccgcgacggc 550	
	teggggetee ceacacetgg ggeetttgee ttecacaceg agtteateea 600	
	cgtccccgtc ggctgcacct gcgtgctgcc ccgttcagtg tgaccgccga 650	
	ggccgtgggg cccctagact ggacacgtgt gctccccaga gggcaccccc 700	
	tatttatgtg tatttattgt tatttatatg cctccccaa cactaccctt 750	
	ggggtetggg catteceegt gtetggagga cageeeeea etgtteteet 800	
	catctccagc ctcagtagtt gggggtagaa ggagctcagc acctcttcca 850	
	gecettaaag etgeagaaaa ggtgteacae ggetgeetgt acettggete 900	
	cctgtcctgc-tcccggcttc_ccttacccta_tcactggcct_caggccccgc 950	

aggctgcctc ttcccaacct ccttggaagt acccctgttt cttaaacaat 1000 tatttaagtg tacgtgtatt attaaactga tgaacacatc cccaaaa 1047

<210> 4

<211> 197

<212> PRT

<213> Homo Sapien

<400> 4

Met Thr Leu Leu Pro Gly Leu Leu Phe Leu Thr Trp Leu His Thr

Cys Leu Ala His His Asp Pro Ser Leu Arg Gly His Pro His Ser

His Gly Thr Pro His Cys Tyr Ser Ala Glu Glu Leu Pro Leu Gly

Gln Ala Pro Pro His Leu Leu Ala Arg Gly Ala Lys Trp Gly Gln

Ala Leu Pro Val Ala Leu Val Ser Ser Leu Glu Ala Ala Ser His

Arg Gly Arg His Glu Arg Pro Ser Ala Thr Thr Gln Cys Pro Val

Leu Arg Pro Glu Glu Val Leu Glu Ala Asp Thr His Gln Arg Ser 95

Ile Ser Pro Trp Arg Tyr Arg Val Asp Thr Asp Glu Asp Arg Tyr 115 110

Pro Gln Lys Leu Ala Phe Ala Glu Cys Leu Cys Arg Gly Cys Ile 130

Asp Ala Arg Thr Gly Arg Glu Thr Ala Ala Leu Asn Ser Val Arg

Leu Leu Gln Ser Leu Leu Val Leu Arg Arg Pro Cys Ser Arg 155

Asp Gly Ser Gly Leu Pro Thr Pro Gly Ala Phe Ala Phe His Thr

Glu Phe Ile His Val Pro Val Gly Cys Thr Cys Val Leu Pro Arg

Ser Val

<210> 5

<211> 1320

<212> DNA

<213> Homo Sapien

<400> 5

ggcttgctga aaataaaatc aggactccta acctgctcca gtcagcctgc 50 ttccacgagg cctgtcagtc agtgcccgac ttgtgactga gtgtgcagtg 100 cccagcatgt accaggtcag tgcagagggc tgcctgaggg ctgtgctgag 150 agggagagga gcagagatgc tgctgagggt ggagggaggc caagctgcca 200 ggtttggggc tgggggccaa gtggagtgag aaactgggat cccaggggga 250 gggtgcagat gagggagcga cccagattag gtgaggacag ttctctcatt 300 agcettttee tacaggtggt tgeattettg geaatggtea tgggaaceca 350 cacctacage cactggeeca getgetgeee cageaaaggg caggaeacet 400 ctgaggaget getgaggtgg ageaetgtge etgtgeetee eetagageet 450 getaggeeca acegeeacee agagteetgt agggeeagtg aagatggaee 500 cctcaacagc agggccatct ccccctggag atatgagttg gacagagact 550 tgaaccggct cccccaggac ctgtaccacg cccgttgcct gtgcccgcac 600 tgcgtcagcc tacagacagg ctcccacatg gacccccggg gcaactcgga 650 gctgctctac cacaaccaga ctgtcttcta caggcggcca tgccatggcg 700 agaagggcac ccacaagggc tactgcctgg agcgcaggct gtaccgtgtt 750 teettagett gtgtgtgtgt geggeeeegt gtgatggget ageeggaeet 800 gctggaggct ggtccctttt tgggaaacct ggagccaggt gtacaaccac 850 ttgccatgaa gggccaggat gcccagatgc ttggcccctg tgaagtgctg 900 tetggageag caggateceg ggacaggatg gggggetttg gggaaaacet 950 gcacttctgc acattttgaa aagagcagct gctgcttagg gccgccggaa 1000 getggtgtee tgteatttte teteaggaaa ggtttteaaa gttetgeeca 1050 tttctggagg ccaccactcc tgtctcttcc tcttttccca tcccctgcta 1100 ccctggccca gcacaggcac tttctagata tttccccctt gctggagaag 1150 aaagagcccc tggttttatt tgtttgttta ctcatcactc agtgagcatc 1200 tactttgggt gcattctagt gtagttacta gtcttttgac atggatgatt 1250 ctgaggagga agctgttatt gaatgtatag agatttatcc aaataaatat 1300 ctttatttaa aaatgaaaaa 1320

<sup>&</sup>lt;210> 6

<sup>&</sup>lt;211> 177

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo-Sapien-

<400> Met 1	6 Arg	Glu	Arg	Pro 5	Arg	Leu	Gly	Glu	Asp 10	Ser	Ser	Leu	Ile	Ser 15
	Phe	Leu	Gln	Val 20	Val	Ala	Phe	Leu	Ala 25	Met	Val	Met	Gly	Thr 30
His	Thr	Tyr	Ser	His 35	Trp	Pro	Ser	Cys	Cys 40	Pro	Ser	Lys	Gly	Gln 45
Asp	Thr	Ser	Glu	Glu 50	Leu	Leu	Arg	Trp	Ser 55	Thr	Val	Pro	Val	Pro 60
Pro	Leu	Glu	Pro	Ala 65	Arg	Pro	Asn	Arg	His 70	Pro	Glu	Ser	Cys	Arg 75
Ala	Ser	Glu	Asp	Gly 80	Pro	Leu	Asn	Ser	Arg 85	Ala	Ile	Ser	Pro	Trp 90
Arg	Tyr	Glu	Leu	Asp 95	Arg	Asp	Leu	Asn	Arg 100	Leu	Pro	Gln	Asp	Leu 105
Tyr	His	Ala	Arg	Cys 110	Leu	Cys	Pro	His	Cys 115	Val	Ser	Leu	Gln	Thr 120
Gly	Ser	His	Met	Asp 125	Pro	Arg	Gly	Asn	Ser 130	Glu	Leu	Leu	Tyr	His 135
Asn	Gln	Thr	Val	Phe	Tyr	Arg	Arg	Pro	Cys 145	His	Gly	Glu	Lys	Gly 150
Thr	His	. Lys	Gly	Tyr 155	Cys	Leu	Glu	Arg	160	J Leu	туг	: Arg	, Val	Ser 165
Leu	ı Ala	a Cys	val	Cys	val	. Arg	J Pro	Arg	Va]	. Met	: Gly	7		

<210> 7

<211> 1754

<212> DNA

<213> Homo Sapien

<400> 7
atgctggtag ccggcttcct gctggcgctg ccgccgagct gggccgcggg 50

cgccccagg gcgggcaggc gccccgcgcg gccgcgggc tgcgcggacc 100

ggccggagga gctactggag cagctgtacg ggcgcctggc ggccgggtg 150

ctcagtgcct tccaccacac gctgcagctg gggccgcgtg agcaggcgc 200

caacgcgagc tgcccggcag ggggcaggcc cggcgaccgc cgcttccggc 250

cgcccaccaa cctgcgcag gtgtcgcct gggcctacag aatctcctac 300

gacccggcga ggtaccccag gtacctgcct gaagcctact gcctgtgcg 350

gggctgcctg-accgggctgt-tcggcgagga-ggacgtgcgc ttccgcagcg 400

cccctgtcta catgcccacc gtcgtcctgc gccgcacccc.cgcctgcgcc 450 ggcggccgtt ccgtctacac cgaggcctac gtcaccatcc ccgtgggctg 500 cacctgcgtc cccgagccgg agaaggacgc agacagcatc aactccagca 550 tcgacaaaca gggcgccaag ctcctgctgg gccccaacga cgcgcccgct 600 ggeeeetgag geeggteetg eeeegggagg teteeeegge eegeateeeg 650 aggegeceaa getggageeg eetggaggge teggteggeg acetetgaag 700 agagtgcacc gagcaaacca agtgccggag caccagcgcc gcctttccat 750 ggagactcgt aagcagcttc atctgacacg ggcatccctg gcttgctttt 800 agctacaagc aagcagcgtg gctggaagct gatgggaaac gacccggcac 850 gggcatcctg tgtgcggccc gcatggaggg tttggaaaag ttcacggagg 900 ctecetgagg ageeteteag ateggetget gegggtgeag ggegtgaete 950 accgctgggt gcttgccaaa gagataggga cgcatatgct ttttaaagca 1000 atctaaaaat aataataagt atagcgacta tatacctact tttaaaatca 1050 actgttttga atagaggcag agctatttta tattatcaaa tgagagctac 1100 tetgttacat ttettaacat ataaacateg ttttttactt ettetggtag 1150 aattttttaa agcataattg gaatccttgg ataaattttg tagctggtac 1200 actetggeet gggtetetga atteageetg teacegatgg etgactgatg 1250 aaatggacac gtctcatctg acccactctt ccttccactg aaggtcttca 1300 cgggcctcca ggtggaccaa agggatgcac aggcggctcg catgccccag 1350 ggccagctaa gagttccaaa gatctcagat ttggttttag tcatgaatac 1400 ataaacagtc tcaaactcgc acaatttttt cccccttttg aaagccactg 1450 gggccaattt gtggttaaga ggtggtgaga taagaagtgg aacgtgacat 1500 ctttgccagt tgtcagaaga atccaagcag gtattggctt agttgtaagg 1550 getttaggat caggetgaat atgaggacaa agtgggeeae gttageatet 1600 gcagagatca atctggaggc ttctgtttct gcattctgcc acgagagcta 1650 ggtccttgat cttttcttta gattgaaagt ctgtctctga acacaattat 1700 ttgtaaaagt tagtagttct tttttaaatc attaaaagag gcttgctgaa 1750 ggat 1754

<210> 8

<211>-202

•	
<212> PRT <213> Homo Sapien	
<pre>&lt;400&gt; 8 Met Leu Val Ala Gly Phe Leu Leu Ala Leu Pro Pro Ser Trp Ala 1 5 10 15</pre>	
Ala Gly Ala Pro Arg Ala Gly Arg Arg Pro Ala Arg Pro Arg Gly 20 25 30	
Cys Ala Asp Arg Pro Glu Glu Leu Leu Glu Gln Leu Tyr Gly Arg 35 40 45	i
Leu Ala Ala Gly Val Leu Ser Ala Phe His His Thr Leu Gln Leu 50 55 60	) }
Gly Pro Arg Glu Gln Ala Arg Asn Ala Ser Cys Pro Ala Gly Gly 65 70	5
Arg Pro Gly Asp Arg Arg Phe Arg Pro Pro Thr Asn Leu Arg Ser 80 85 90	<i>-</i> )
Val Ser Pro Trp Ala Tyr Arg Ile Ser Tyr Asp Pro Ala Arg Tyr 95 100	<u>-</u> 5
Pro Arg Tyr Leu Pro Glu Ala Tyr Cys Leu Cys Arg Gly Cys Leu 110 115 120	-
Thr Gly Leu Phe Gly Glu Glu Asp Val Arg Phe Arg Ser Ala Pro 125 130	5 5
Val Tyr Met Pro Thr Val Val Leu Arg Arg Thr Pro Ala Cys Ala 140 145	a 0
Gly Gly Arg Ser Val Tyr Thr Glu Ala Tyr Val Thr Ile Pro Va 155 160 16	1 5
Gly Cys Thr Cys Val Pro Glu Pro Glu Lys Asp Ala Asp Ser II 170 175 18	e 0
Asn Ser Ser Ile Asp Lys Gln Gly Ala Lys Leu Leu Gly Pr 185 190 19	5
Asn Asp Ala Pro Ala Gly Pro 200	
<210> 9 <211> 559 <212> DNA <213> Homo Sapien	
<400> 9 caactgcacc tcggttctat cgatagccac cagcgcaaca tgacagtgaa 50	
gaccetgeat ggeceageea tggteaagta ettgetgetg tegatattgg 100	)

ggcttgcctt tctgagtgag gcggcagctc ggaaaatccc caaagtagga 150

catactttt tocaaaagce tgagagttge cegeetgtge caggaggtag 200 tatgaagett gacattggca teateaatga aaaecagege gtttecatgt 250 caegtaacat egagageege teeaecteee eetggaatta caetgteaet 300 tgggaceeca aecggtaeee eteggaagtt gtacaggeee agtgtaggaa 350 ettgggetge ateaatgete aaggaaagga agacatetee atgaatteeg 400 teegteteet teeagttgga gaaggtgetg gtgaetgtt getgeaeetg 500 egteaeeeet gteateeae atgtgeagta agaggtgeat ateeaeteag 550 etgaagaag 559

<210> 10 <211> 163 <212> PRT

<213> Homo Sapien

Ile Ile Asn Glu Asn Gln Arg Val Ser Met Ser Arg Asn Ile Glu
65 70 75

Ser Arg Ser Thr Ser Pro Trp Asn Tyr Thr Val Thr Trp Asp Pro 80 85 90

Asn Arg Tyr Pro Ser Glu Val Val Gln Ala Gln Cys Arg Asn Leu 95 100 105

Gly Cys Ile Asn Ala Gln Gly Lys Glu Asp Ile Ser Met Asn Ser 110 115

Val Pro Ile Gln Gln Glu Thr Leu Val Val Arg Arg Lys His Gln 125 130 135

Gly Cys Ser Val Ser Phe Gln Leu Glu Lys Val Leu Val Thr Val

Gly Cys Thr Cys Val Thr Pro Val Ile His His Val Gln 155 160

<2.1-0->--1-1

<211> 1515 <212> DNA

<213> Homo Sapien

<400> 11 ceggegatgt egetegtget getaageetg geegegetgt geaggagege 50 cgtaccecga gagccgaccg ttcaatgtgg ctctgaaact gggccatctc 100 cagagtggat gctacaacat gatctaatcc ccggagactt gagggacctc 150 cgagtagaac ctgttacaac tagtgttgca acaggggact attcaatttt 200 gatgaatgta agctgggtac tccgggcaga tgccagcatc cgcttgttga 250 aggccaccaa gatttgtgtg acgggcaaaa gcaacttcca gtcctacagc 300 tgtgtgaggt gcaattacac agaggccttc cagactcaga ccagaccctc 350 tggtggtaaa tggacatttt cctacatcgg cttccctgta gagctgaaca 400 cagtctattt cattggggcc cataatattc ctaatgcaaa tatgaatgaa 450 gatggccctt ccatgtctgt gaatttcacc tcaccaggct gcctagacca 500 cataatgaaa tataaaaaaa agtgtgtcaa ggccggaagc ctgtgggatc 550 cgaacatcac tgcttgtaag aagaatgagg agacagtaga agtgaacttc 600 acaaccactc ccctgggaaa cagatacatg gctcttatcc aacacagcac 650 tatcatcggg ttttctcagg tgtttgagcc acaccagaag aaacaaacgc 700 gagetteagt ggtgatteea gtgaetgggg atagtgaagg tgetaeggtg 750 cagctgactc catattttcc tacttgtggc agcgactgca tccgacataa 800 aggaacagtt gtgctctgcc cacaaacagg cgtccctttc cctctggata 850 acaacaaaag caagccggga ggctggctgc ctctcctcct gctgtctctg 900 ctggtggcca catgggtgct ggtggcaggg atctatctaa tgtggaggca 950 cgaaaggatc aagaagactt ccttttctac caccacacta ctgcccccca 1000 ttaaggttct tgtggtttac ccatctgaaa tatgtttcca tcacacaatt 1050 tgttacttca ctgaatttct tcaaaaccat tgcagaagtg aggtcatcct 1100 tgaaaagtgg cagaaaaaga aaatagcaga gatgggtcca gtgcagtggc 1150 ttgccactca aaagaaggca gcagacaaag tcgtcttcct tctttccaat 1200 gacgtcaaca gtgtgtgcga tggtacctgt ggcaagagcg agggcagtcc 1250 cagtgagaac teteaagaec tetteeecet tgeetttaac ettttetgea 1300 gtgatctaag aagccagatt\_catctgcaca\_aatacgtggt\_ggtctacttt 1350 agagagattg atacaaaaga cgattacaat gctctcagtg tctgccccaa 1400 gtaccacctc atgaaggatg ccactgcttt ctgtgcagaa cttctccatg 1450 tcaagcagca ggtgtcagca ggaaaaagat cacaagcctg ccacgatggc 1500 tgctgctcct tgtag 1515

<210> 12

<211> 502

<212> PRT

<213> Homo Sapien

<400> 12

Me	t 1			Val	_		Ser	Leu	Ala	Ala 10	Leu	Cys	Arg	Ser	15
۷a	.1	Pro	Arg	Glu	Pro 20	Thr	Val	Gln	Cys	Gly 25	Ser	Glu	Thr	Gly	Pro 30

Ser Pro Glu Trp Met Leu Gln His Asp Leu Ile Pro Gly Asp Leu

Arg Asp Leu Arg Val Glu Pro Val Thr Thr Ser Val Ala Thr Gly
50 55 60

Asp Tyr Ser Ile Leu Met Asn Val Ser Trp Val Leu Arg Ala Asp 65 70 75

Ala Ser Ile Arg Leu Leu Lys Ala Thr Lys Ile Cys Val Thr Gly 80 85 90

Lys Ser Asn Phe Gln Ser Tyr Ser Cys Val Arg Cys Asn Tyr Thr 95 100 105

Glu Ala Phe Gln Thr Gln Thr Arg Pro Ser Gly Gly Lys Trp Thr

Phe Ser Tyr Ile Gly Phe Pro Val Glu Leu Asn Thr Val Tyr Phe 125 130 135

Ile Gly Ala His Asn Ile Pro Asn Ala Asn Met Asn Glu Asp Gly 140 145

Pro Ser Met Ser Val Asn Phe Thr Ser Pro Gly Cys Leu Asp His 155 160 165

Ile Met Lys Tyr Lys Lys Lys Cys Val Lys Ala Gly Ser Leu Trp 170 175 180

Asp Pro Asn Ile Thr Ala Cys Lys Lys Asn Glu Glu Thr Val Glu 185 190

Val Asn Phe Thr Thr Thr Pro Leu Gly Asn Arg Tyr Met Ala Leu 200 205 210

Ile Gln His Ser Thr Ile Ile Gly Phe Ser Gln Val Phe Glu Pro215220225

His	Gln	Lys	Lys	Gln 230	Thr	Arg	Ala	Ser	Val 235	Val	Ile	Pro	Val	Thr 240
Gly	Asp	Ser	Glu	Gly 245	Ala	Thr	Val	Gln	Leu 250	Thr	Pro	Tyr	Phe	Pro 255
Thr	Cys	Gly	Ser	Asp 260	Cys	Ile	Arg	His	Lys 265	Gly	Thr	Val	Val	Leu 270
Cys	Pro	Gln	Thr	Gly 275	Val	Pro	Phe	Pro	Leu 280	Asp	Asn	Asn	Lys	Ser 285
Lys	Pro	Gly	Gly	Trp 290	Leu	Pro	Leu	Leu	Leu 295	Leu	Ser	Leu	Leu	Val 300
Ala	Thr	Trp	Val	Leu 305	Val	Ala	Gly	Ile	Tyr 310	Leu	Met	Trp	Arg	His 315
Glu	Arg	Ile	Lys	Lys 320	Thr	Ser	Phe	Ser	Thr 325	Thr	Thr	Leu	Leu	Pro 330
Pro	Ile	Lys	Val	Leu 335	Val	Val	Tyr	Pro	Ser 340	Glu	Ile	Cys	Phe	His 345
His	Thr	Ile	Cys	Tyr 350	Phe	Thr	Glu	Phe	Leu 355	Gln	Asn	His	Cys	Arg 360
Ser	Glu	Val	Ile	Leu 365	Glu	Lys	Trp	Gln	Lys 370	Lys	Lys	Ile	Ala	Glu 375
Met	Gly	Pro	Val	Gln 380	Trp	Leu	Ala	Thr	Gln 385	Lys	Lys	Ala	Ala	Asp 390
Lys	Val	Val	Phe	Leu 395	Leu	Ser	Asn	Asp	Val 400	Asn	Ser	Val	Cys	Asp 405
_				Lys 410					415					420
Asp	Leu	Phe	Pro	Leu 425	Ala	Phe	Asn	Leu	Phe 430	Cys	Ser	Asp	Leu	Arg 435
Ser	Gln	Ile	His	Leu 440	His	Lys	Tyr	Val	Val 445	Val	Tyr	Phe	Arg	Glu 450
Ile	Asp	Thr	Lys	Asp 455	Asp	Tyr	Asn	Ala	Leu 460	Ser	Val	Cys	Pro	Lys 465
Tyr	His	Leu	Met	Lys 470	Asp	Ala	Thr	Ala	Phe 475	Cys	Ala	Glu	Leu	Leu 480
His	Val	Lys	Gln	Gln 485	Val	Ser	Ala	Gly	Lys 490	Arg	Ser	Gln	Ala	Cys 495
His	Asp	Gly	Cys	Cys 500	Ser	Leu								
-210	. 12													

<211> 2380

<212> DNA

<213> Homo Sapien

<400> 13 acactggcca aacaaaaacg aaagcactcc gtgctggaag taggaggaga 50 gtcaggactc ccaggacaga gagtgcacaa actacccagc acagccccct 100 ccgcccctc tggaggctga agagggattc cagcccctgc cacccacaga 150 cacgggctga ctggggtgtc tgccccctt gggggggggc agcacagggc 200 ctcaggcctg ggtgccacct ggcacctaga agatgcctgt gccctggttc 250 ttgctgtcct tggcactggg ccgaagccca gtggtccttt ctctggagag 300 gcttgtgggg cctcaggacg ctacccactg ctctccgggc ctctcctgcc 350 gcctctggga cagtgacata ctctgcctgc ctggggacat cgtgcctgct 400 ccgggccccg tgctggcgcc tacgcacctg cagacagagc tggtgctgag 450 gtgccagaag gagaccgact gtgacctctg tctgcgtgtg gctgtccact 500 tggccgtgca tgggcactgg gaagagcctg aagatgagga aaagtttgga 550 ggagcagctg actcaggggt ggaggagcct aggaatgcct ctctccaggc 600 ccaagtcgtg ctctccttcc aggcctaccc tactgcccgc tgcgtcctgc 650 tggaggtgca agtgcctgct gcccttgtgc agtttggtca gtctgtgggc 700 tctgtggtat atgactgctt cgaggctgcc ctagggagtg aggtacgaat 750 ctggtcctat actcagccca ggtacgagaa ggaactcaac cacacagc 800 agetgeetge cetgeeetgg etcaaegtgt cageagatgg tgacaaegtg 850 catctggttc tgaatgtctc tgaggagcag cacttcggcc tctccctgta 900 ctggaatcag gtccagggcc ccccaaaacc ccggtggcac aaaaacctga 950 ctggaccgca gatcattacc ttgaaccaca cagacctggt tccctgcctc 1000 tgtattcagg tgtggcctct ggaacctgac tccgttagga cgaacatctg 1050 ccccttcagg gaggaccccc gcgcacacca gaacctctgg caagccgccc 1100 gactgcgact gctgaccctg cagagctggc tgctggacgc accgtgctcg 1150 ctgcccgcag aagcggcact gtgctggcgg gctccgggtg gggacccctg 1200 ccagccactg gtcccaccgc tttcctggga gaacgtcact gtggacaagg 1250 ttctcgagtt cccattgctg aaaggccacc ctaacctctg tgttcaggtg 1300 aacagetegg agaagetgea getgeaggag tgettgtggg etgaeteeet 1350

ggggcctctc aaagacgatg tgctactgtt ggagacacga ggcccccagg 1400 acaacagatc cctctgtgcc ttggaaccca gtggctgtac ttcactaccc 1450 agcaaagcct ccacgagggc agctcgcctt ggagagtact tactacaaga 1500 cctgcagtca ggccagtgtc tgcagctatg ggacgatgac ttgggagcgc 1550 tatgggcctg ccccatggac aaatacatcc acaagcgctg ggccctcgtg 1600 tggctggcct gcctactctt tgccgctgcg ctttccctca tcctccttct 1650 caaaaaggat cacgcgaaag ggtggctgag gctcttgaaa caggacgtcc 1700 gctcgggggc ggccgccagg ggccgcgcgg ctctgctcct ctactcagcc 1750 gatgactcgg gtttcgagcg cctggtgggc gccctggcgt cggccctgtg 1800 ccagctgccg ctgcgcgtgg ccgtagacct gtggagccgt cgtgaactga 1850 gcgcgcaggg gcccgtggct tggtttcacg cgcagcggcg ccagaccctg 1900 caggagggg gcgtggtggt cttgctcttc tctcccggtg cggtggcgct 1950 gtgcagcgag tggctacagg atggggtgtc cgggcccggg gcgcacggcc 2000 cgcacgacgc cttccgcgcc tcgctcagct gcgtgctgcc cgacttcttg 2050 cagggccggg cgcccggcag ctacgtgggg gcctgcttcg acaggctgct 2100 ccacccggac gccgtacccg cccttttccg caccgtgccc gtcttcacac 2150 tgccctccca actgccagac ttcctggggg ccctgcagca gcctcgcgcc 2200 ccgcgttccg ggcggctcca agagagagcg gagcaagtgt cccgggccct 2250 tcagccagcc ctggatagct acttccatcc cccggggact cccgcgccgg 2300 gacgcggggt gggaccaggg gcgggacctg gggcggggga cgggacttaa 2350 ataaaggcag acgctgtttt tctaaaaaaa 2380

## <400> 14

Met Pro Val Pro Trp Phe Leu Leu Ser Leu Ala Leu Gly Arg Ser 1 5 10 15

Pro Val Val Leu Ser Leu Glu Arg Leu Val Gly Pro Gln Asp Ala 20 25 30

Thr His Cys Ser Pro Gly Leu Ser Cys Arg Leu Trp Asp Ser Asp 35 40 45

Ile Leu Cys Leu Pro Gly Asp Ile Val Pro Ala Pro Gly Pro Val

<sup>&</sup>lt;210> 14

<sup>&</sup>lt;211> 705

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

		•		50					55					60
Leu	Ala	Pro	Thr	His 65	Leu	Gln	Thr	Glu	Leu 70	Val	Leu	Arg	Cys	Gln 75
Lys	Glu	Thr	Asp	Cys 80	Asp	Leu	Cys	Leu	Arg 85	Val	Ala	Val	His	Leu 90
Ala	Val	His	Gly	His 95	Trp	Glu	Glu	Pro	Glu 100	Asp	Glu	Glu	Lys	Phe 105
Gly	Gly	Ala	Ala	Asp 110	Ser	Gly	Val	Glu	Glu 115	Pro	Arg	Asn	Ala	Ser 120
Leu	Gln	Ala	Gln	Val 125	Val	Leu	Ser	Phe	Gln 130	Ala	Tyr	Pro	Thr	Ala 135
Arg	Cys	Val	Leu	Leu 140	Glu	Val	Gln	Val	Pro 145	Ala	Ala	Leu	Val	Gln 150
Phe	Gly	Gln	Ser	Val 155	Gly	Ser	Val	Val	Tyr 160	Asp	Cys	Phe	Glu	Ala 165
Ala	Leu	Gly	Ser	Glu 170	Val	Arg	Ile	Trp	Ser 175	Tyr	Thr	Gln	Pro	Arg 180
Tyr	Glu	Lys	Glu	Leu 185	Asn	His	Thr	Gln	Gln 190	Leu	Pro	Ala	Leu	Pro 195
Trp	Leu	Asn	Val	Ser 200	Ala	Asp	Gly	Asp	Asn 205	Val	His	Leu	Val	Leu 210
Asn	Val	Ser	Glu	Glu 215	Gln	His	Phe	Gly	Leu 220	Ser	Leu	Tyr	Trp	Asn 225
Gln	Val	Gln	Gly	Pro 230	Pro	Lys	Pro	Arg	Trp 235	His	Lys	Asn	Leu	Thr 240
Gly	Pro	Gln	Ile	Ile 245	Thr	Leu	Asn	His	Thr 250	Asp	Leu	Val	Pro	Cys 255
Leu	Cys	Ile	Gln	Val 260	Trp	Pro	Leu	Glu	Pro 265	Asp	Ser	Val	Arg	Thr 270
Asn	Ile	Cys	Pro	Phe 275	Arg	Glu	Asp	Pro	Arg 280	Ala	His	Gln	Asn	Leu 285
Trp	Gln	Ala	Ala	Arg 290	Leu	Arg	Leu	Leu	Thr 295	Leu	Gln	Ser	Trp	Leu 300
Leu	Asp	Ala	Pro	Cys 305	Ser	Leu	Pro	Ala	Glu 310	Ala	Ala	Leu	Cys	Trp 315
Arg	Ala	Pro	Gly	Gly 320	Asp	Pro	Cys	Gln	Pro 325	Leu	Val	Pro	Pro	Leu 330
Ser	Trp	Glu	Asn	Val 335	Thr	Val	Asp	Lys	Val 340	Leu	Glu	Phe	Pro	Leu 345

Leu         Gln         Leu         Gln         Gln         Cys         Leu         Trp         Asp         Asp         Leu         Gln         Pro         Saro         Asp         Asp         Asp         Asp         Asp         Leu         Leu         Leu         Glu         Trp         Ser         Gly         Pro         Gly         Pro         Asp           Asn         Arg         Leu         Gly         Ala         Leu         Glu         Pro         Ser         Gly         Glu         Trp         Asp	Leu	Lys	Gly	His	Pro 350	Asn	Leu	Cys	Val	Gln 355	Val	Asn	Ser	Ser	Glu 360
Asn       Arg       Ser       Leu       Cys Ala Leu       Glu Pro Ser Gly Cys Thr Ser Leu A05         Pro       Ser Lys Ala Ser Leu Gln Ser Gly Glu Pro Ser Lys Ala Ser Gly Glu Pro Ser Lys Ala Ser Gly Glu Glu Tyr Leu A15       Leu Gln Asp Leu Gln Ser Gly Gln Cys Leu Gln Leu Trp Asp Asp A35         Asp Leu Gly Ala Leu Gln Ser Gly Glu Pro Ser Gly Glu Pro Ser Gly Glu Pro Ser Gly Glu Pro Ser Gly Glu Fry Glo Ser Gly Gly Gly Gly Glu Fry Glo Ser Gly Glu Fry Glo Ser Gly Gly Gly Gly Fry Glo Ser Gly	Lys	Leu	Gln	Leu		Glu	Суѕ	Leu	Trp		Asp	Ser	Leu	Gly	
19	Leu	Lys	Asp	Asp		Leu	Leu	Leu	Glu		Arg	Gly	Pro	Gln	
Leu Gln Asp Leu Gln Asp Leu Gln Ser Gly Gln Cys Leu Gln Leu Trp Asp Asp A35           Asp Leu Gly Ala Leu Trp Asp Asp Lys Tyr Ile His Asp Lys Arg Trp Ala Leu A40         Trp Ala Cys Pro Met A45         Asp Leu Gly Ala Leu A10         Trp Leu A40         Trp Leu Lug A45         Leu Lug Lug A45         Leu Lug Lug A45         Leu Lug Lug A40         Leu Lug Lug A45         Leu Lug Lug A45         Leu Lug Lug A45         Lug A45 <td>Asn</td> <td>Arg</td> <td>Ser</td> <td>Leu</td> <td>-</td> <td>Ala</td> <td>Leu</td> <td>Glu</td> <td>Pro</td> <td></td> <td>Gly</td> <td>Cys</td> <td>Thr</td> <td>Ser</td> <td></td>	Asn	Arg	Ser	Leu	-	Ala	Leu	Glu	Pro		Gly	Cys	Thr	Ser	
Asp         Leu         Gly         Ala         Leu degree         Ala         Cys         Pro degree         Met degree         Asp Lys         Tyr         Ile degree         His degree           Lys         Arg         Trp         Ala         Leu degree         Val         Trp         Leu degree         Ala         Cys         Leu degree         Leu degree         Ala         Leu degree         Ala         Ala         Ala         Ala         Leu degree         Ala	Pro	Ser	Lys	Ala		Thr	Arg	Ala	Ala		Leu	Gly	Glu	Tyr	
Lys Arg Trp Ala       Leu Val Trp Leu Ala Cys Leu Leu Phe Ala Ala 465         Ala Leu Ser Leu Ile Leu Lys Blan Asp Leu Lys Lys Asp His Ala Lys Gly 470         Trp Leu Arg Leu Leu Lys Blan Asp Leu Leu Lys Blan Asp Blan Asp Asp Blan Asp Asp Blan Ala Ala 490         Arg Gly Arg Ala Ala Ala Leu Leu Leu Leu Tyr Ser Ala Asp Asp Asp Ser Gly 510         Phe Glu Arg Leu Val Gly Ala Leu Leu Leu Trp Ser Ala Leu Cys Gln Leu 525         Pro Leu Arg Val Ala Val Asp Leu Trp Ser Arg Arg Glu Leu Ser 530         Ala Gln Gly Pro Val Ala Trp Phe His Ala Gln Arg Arg Gln Thr 545         Leu Gln Glu Gly Pro Val Ala Val Val Val Leu Leu Phe Ser Pro Gly Ala 560         Val Ala Leu Cys Ser Glu Trp Leu Gln Asp Gly Val Ser Gly 570         Val Ala His Gly Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 605         Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Val Val Gly Arg Ala Pro Gly Ala Cys Bro Got         Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Val Val Gly Ala Gro Gly Pro Gly Ala Gro Gro Gly Ala Gro Gly Ala Gro Gly Ala Gro Gly Ala Gro Gro Gly Ala Gro Gly Ala Gro	Leu	Gln	Asp	Leu		Ser	Gly	Gln	Cys		Gln	Leu	Trp	Asp	
Ala       Leu       Ser       Leu       Leu       Leu       Leu       Leu       Lys       Lys       Asp       His       Ala       Lys       Gly         Trp       Leu       Arg       Leu       Leus       Lys       Gln       Asp       Val       Arg       Ala	Asp	Leu	Gly	Ala		Trp	Ala	Cys	Pro		Asp	Lys	Tyr	Ile	
Trp         Leu         Arg         Leu         Leu         Leu         Lys         Gln         Asp         Val         Arg         Ser         Gly         Ala         Ala <td>Lys</td> <td>Arg</td> <td>Trp</td> <td>Ala</td> <td></td> <td>Val</td> <td>Trp</td> <td>Leu</td> <td>Ala</td> <td></td> <td>Leu</td> <td>Leu</td> <td>Phe</td> <td>Ala</td> <td></td>	Lys	Arg	Trp	Ala		Val	Trp	Leu	Ala		Leu	Leu	Phe	Ala	
Arg       Gly       Arg       Ala       Ala something states and states and states and states and states are states and states and states are states and states are	Ala	Leu	Ser	Leu		Leu	Leu	Leu	Lys		Asp	His	Ala	Lys	
Phe         Glu         Arg         Leu         Val 515         Gly         Ala         Leu         Ala         Ser Ala         Leu         Cys         Glu         Leu 525           Pro         Leu         Arg         Val         Ala         Val         Asp         Leu         Trp         Ser Arg         Arg         Glu         Leu         Ser 540           Ala         Gln         Gly         Val         Ala         Trp         Phe         His         Ala         Arg         Arg         Glu         Leu         Ser 540           Ala         Gln         Gly         Val         Ala         Trp         Phe         His         Ala         Arg         Arg         Arg         Gln         Trp 555           Leu         Gln         Gly         Gly         Val         Val         Val         Leu         Leu         Leu         Phe         Ser F85         Phe         Ser F85         Pro         Gly         Arg         Ala         Pro         Ser Gly	Trp	Leu	Arg	Leu		Lys	Gln	Asp	Val		Ser	Gly	Ala	Ala	
Pro       Leu       Arg       Val       Ala Sol	Arg	Gly	Arg	Ala		Leu	Leu	Leu	Tyr		Ala	Asp	Asp	Ser	
Ala       Gln       Gly       Pro       Val 545       Ala       Trp       Phe       His 550       Gln       Arg       Arg       Gln       Thr 555         Leu       Gln       Gly       Gly       Val       Val       Leu       Leu       Phe       Ser       Pro       Gly       Ala         Val       Ala       Leu       Cys       Ser       Glu       Trp       Leu       Gln       Asp       Gly       Val       Ser       Gly       Pro       585         Gly       Ala       His       Gly       Pro       His       Asp       Ala       Phe       Arg       Ala       Pro       Gly       Ser       Cys       Goo       Cys       Goo       Cys       Goo       Arg       Ala       Pro       Gly       Ser       Tyr       Val       Gly       Ala       Gly       Ala       Cys       Ala       Gly       Pro       Ala       Gly       Pro       Ala       Gly       Ala       Gly       Ala       Cys       Ala       Gly       Ala       Gly       Ala       Fro       Ala       Gly       Ala       Gly       Ala       Gly       Ala       Gly       Ala       Gly	Phe	Glu	Arg	Leu		Gly	Ala	Leu	Ala		Ala	Leu	Cys	Gln	
Ala Gln Gly Pro Val Ala Trp Phe His Ala Gln Arg Arg Gln Thr 555  Leu Gln Glu Gly Gly Val Val Val Leu Leu Phe Ser Pro Gly Ala 570  Val Ala Leu Cys Ser Glu Trp Leu Gln Asp Gly Val Ser Gly Pro 585  Gly Ala His Gly Pro His Asp Ala Phe Arg Ala Ser Leu Ser Cys 600  Val Leu Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 615  Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 630	Pro	Leu	Arg	Val		Val	Asp	Leu	Trp		Arg	Arg	Glu	Leu	
Leu Gln Glu Gly Gly Val Val Val Leu Leu Gln Ser Pro Gly Ala 570  Val Ala Leu Cys Ser Glu Trp Leu Gln Asp Gly Val Ser Gly Pro 585  Gly Ala His Gly Pro His Asp Ala Phe Arg Ala Ser Leu Ser Cys 600  Val Leu Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 615  Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 630	_	_				_					<u>.</u>		_		
Val Ala Leu Cys Ser Glu Trp Leu Gln Asp Gly Val Ser Gly Pro 585  Gly Ala His Gly Pro His Asp Ala Phe Arg Ala Ser Leu Ser Cys 600  Val Leu Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 615  Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 630					545					550					555
Gly Ala His Gly Pro His Asp Ala Phe Arg Ala Ser Leu Ser Cys 600  Val Leu Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 615  Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 630	Leu	Gln	Glu	Gly	_	Val	Val	Val	Leu		Phe	Ser	Pro	Gly	
Val Leu Pro Asp Phe Leu Gln Gly Arg Ala Pro Gly Ser Tyr Val 615 Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 620	Val	Ala	Leu	Cys		Glu	Trp	Leu	Gln		Gly	Val	Ser	Gly	
Gly Ala Cys Phe Asp Arg Leu Leu His Pro Asp Ala Val Pro Ala 620 630	Gly	Ala	His	Gly		His	Asp	Ala	Phe		Ala	Ser	Leu	Ser	
620 625 630	Val	Leu	Pro	Asp		Leu	Gln	Gly	Arg		Pro	Gly	Ser	Tyr	
Leu Phe Arg Thr Val Pro Val Phe Thr Leu Pro Ser Gln Leu Pro	Gly	Ala	Cys	Phe		Arg	Leu	Leu	His		Asp	Ala	Val	Pro	
	Leu	Phe	Arg	Thr	Val	Pro	Val	Phe	Thr	Leu	Pro	Ser	Gln	Leu	Pro

640 645 635 Asp Phe Leu Gly Ala Leu Gln Gln Pro Arg Ala Pro Arg Ser Gly Arg Leu Gln Glu Arg Ala Glu Gln Val Ser Arg Ala Leu Gln Pro 675 670 Ala Leu Asp Ser Tyr Phe His Pro Pro Gly Thr Pro Ala Pro Gly Arg Gly Val Gly Pro Gly Ala Gly Pro Gly Ala Gly Asp Gly Thr <210> 15 <211> 2138 <212> DNA <213> Homo Sapien <400> 15 cgagggetee tgetggtaet gtgttegetg etgeacagea aggeeetgee 50 acceaectte aggecatgea gecatgttee gggageecta attgeaeaga 100 agcccatggg gagctccaga ctggcagccc tgctcctgcc tctcctcctc 150 atagtcatcg acctetetga etetgetggg attggettte gecacetgee 200 ccactggaac acccgctgtc ctctggcctc ccacacggat gacagtttca 250 ctggaaqttc tqcctatatc ccttgccgca cctggtgggc cctcttctcc 300 acaaaqcctt qqtqtqtqcq agtctgqcac tqttcccqct qtttgtgcca 350 gcatctqctq tcaggtggct caggtcttca acggggcctc ttccacctcc 400 tggtgcagaa atccaaaaag tcttccacat tcaagttcta taggagacac 450

aagatgccag cacctgctca gaggaagctg ctgcctcgtc gtcacctgtc 500

tgagaagagc catcacattt ccatcccctc cccagacatc tcccacaagg 550

gacttcgctc taaaaggacc caaccttcgg atccagagac atgggaaagt 600

cttcccagat tggactcaca aaggcatgga ggacccgagt tctcctttga 650

tttgctgcct gaggcccggg ctattcgggt gaccatatct tcaggccctg 700

aggtcagcgt gcgtctttgt caccagtggg cactggagtg tgaagagctg 750

agcagtccct atgatgtcca gaaaattgtg tctgggggcc acactgtaga 800

gctgccttat gaattccttc tgccctgtct gtgcatagag gcatcctacc 850

tgcaagagga cactgtgagg cgcaaaaaat gtcccttcca gagctggcca 900

gaagcctatg gctcggactt ctggaagtca gtgcacttca ctgactacag 950

ccagcacact. cagatggtca tggccctgac actccgctgc ccactgaagc 1000 tggaagctgc cctctgccag aggcacgact ggcataccct ttgcaaagac 1050 ctcccgaatg ccacggctcg agagtcagat gggtggtatg ttttggagaa 1100 ggtggacctg caccccagc tctgcttcaa gttctctttt ggaaacagca 1150 gccatgttga atgcccccac cagactgggt ctctcacatc ctggaatgta 1200 agcatggata cccaagccca gcagctgatt cttcacttct cctcaagaat 1250 gcatgccacc ttcagtgctg cctggagcct cccaggcttg gggcaggaca 1300 ctttggtgcc ccccgtgtac actgtcagcc aggcccgggg ctcaagccca 1350 gtgtcactag acctcatcat tecetteetg aggecagggt getgtgteet 1400 ggtgtggcgg tcagatgtcc agtttgcctg gaagcacctc ttgtgtccag 1450 atgtetetta cagacacetg gggetettga teetggeact getggeeete 1500 ctcaccctac tgggtgttgt tctggccctc acctgccggc gcccacagtc 1550 aggcccgggc ccagcgcggc cagtgctcct cctgcacgcg gcggactcgg 1600 aggcgcagcg gcgcctggtg ggagcgctgg ctgaactgct acgggcagcg 1650 ctgggcggcg ggcgcgacgt gatcgtggac ctgtgggagg ggaggcacgt 1700 ggcgcgcgtg ggcccgctgc cgtggctctg ggcggcgcgg acgcgcgtag 1750 cgcgggagca gggcactgtg ctgctgctgt ggagcggcgc cgaccttcgc 1800 ceggteageg geceegacee eegegeegeg eeeetgeteg eeetgeteea 1850 cgctgccccg cgcccgctgc tgctgctcgc ttacttcagt cgcctctgcg 1900 ccaagggcga catcccccg ccgctgcgcg ccctgccgcg ctaccgcctg 1950 ctgcgcgacc tgccgcgtct gctgcgggcg ctggacgcgc ggcctttcgc 2000 agaggccacc agctggggcc gccttggggc gcggcagcgc aggcagagcc 2050 gcctagagct gtgcagccgg cttgaacgag aggccgcccg acttgcagac 2100 ctaggttgag cagageteca cegeagtece gggtgtet 2138

<sup>&</sup>lt;210> 16

<sup>&</sup>lt;211> 667

<sup>&</sup>lt;212> PRT

<sup>&</sup>lt;213> Homo Sapien

<sup>&</sup>lt;400> 16

Met Gly Ser Ser Arg Leu Ala Ala Leu Leu Leu Pro Leu Leu Leu 1 5 10 15

Ile Val Ile Asp Leu Ser Asp Ser Ala Gly Ile Gly Phe Arg His

		•		20					25		,			30
Leu	Pro	His	Trp		Thr	Arg	Cys	Pro	Leu 40	Ala	Ser	His	Thr	Asp 45
Asp	Ser	Phe	Thr		Ser	Ser	Ala	Tyr		Pro	Cys	Arg	Thr	Trp 60
Trp	Ala	Leu	Phe		Thr	Lys	Pro	Trp	Cys 70	Val	Arg	Val	Trp	His 75
Cys	Ser	Arg	Cys		Cys	Gln	His	Leu	Leu 85	Ser	Gly	Gly	Ser	Gly 90
Leu	Gln	Arg	Gly	Leu 95	Phe	His	Leu	Leu	Val 100	Gln	Lys	Ser	Lys	Lys 105
Ser	Ser	Thr	Phe	Lys 110	Phe	Tyr	Arg	Arg	His 115	Lys	Met	Pro	Ala	Pro 120
Ala	Gln	Arg	Lys	Leu 125	Leu	Pro	Arg	Arg	His 130	Leu	Ser	Glu	Lys	Ser 135
His	His	Ile	Ser	Ile 140	Pro	Ser	Pro	Asp	Ile 145	Ser	His	Lys	Gly	Leu 150
Arg	Ser	Lys	Arg	Thr 155	Gln	Pro	Ser	Asp	Pro 160	Glu	Thr	Trp	Glu	Ser 165
Leu	Pro	Arg	Leu	Asp 170	Ser	Gln	Arg	His	Gly 175	Gly	Pro	Glu	Phe	Ser 180
Phe	Asp	Leu	Leu	Pro 185	Glu	Ala	Arg	Ala	Ile 190	Arg	Val	Thr	Ile	Ser 195
Ser	Gly	Pro	Glu	Val 200	Ser	Val	Arg	Leu	Cys 205	His	Gln	Trp	Ala	Leu 210
Glu	Cys	Glu	Glu	Leu 215	Ser	Ser	Pro	Tyr	Asp 220	Val	Gln	Lys	Ile	Val 225
Ser	Gly	· Gly	His	Thr 230	Val	Glu	Leu	Pro	Tyr 235	Glu	Phe	Leu	Leu	Pro 240
Cys	Leu	Cys	Ile	Glu 245		Ser	Tyr	Leu	Gln 250	Glu	. Asp	Thr	Val	Arg 255
Arg	Lys	Lys	Cys	Pro 260		Gln	. Ser	Trp	Pro 265	Glu	Ala	Туг	Gly	Ser 270
Asp	Phe	Trp	. Lys	Ser 275		His	Phe	Thr	Asp 280	туг	Ser	Glr	n His	Thr 285
Gln	Met	: Val	. Met	Ala 290		Thr	Leu	Arg	Cys 295	Pro	Leu	Lys	: Leu	Glu 300
Ala	Ala	Leu	Cys	Gln 305		His	Asp	Trp	310	Thr	Leu	. Суя	Lys	315

Glu Lys Val Asp Leu His Pro Gln Leu Cys Phe Lys Phe Ser Phe 345  Gly Asn Ser Ser His Val Glu Cys Pro His Gln Thr Gly Ser Leu 350  Thr Ser Trp Asn Val Ser Met Asp Thr Gln Ala Gln Gln Leu Ile 370  Leu His Phe Ser Ser Arg Met His Ala Thr 385  Ser Leu Pro Gly Leu Gly Gln Asp Thr Leu Val Pro Pro Val Trp 395  Thr Val Ser Gln Ala Arg Gly Ser Ser Pro Val Ser Leu Asp Leu 415  Ser Asp Val Gln Phe Ala Trp Lys His Leu Leu Cys Pro Asp Val 485  Leu Thr Leu Leu Gly Val Val Leu Ala Leu Trr Cys Arg Arg Pro 480  Gln Ser Gly Pro Gly Pro Ala Arg Pro Val Leu Leu His Ala Ala App Chr Asp Ser Gly Asp Ser Gly Gly Gly Asp Asp Val Ile Val Asp 525  Leu Trp Glu Gly Arg His Val Ala Arg Val Ala Arg Gly Gly Gly Arg Arg Val Ile Val Asp 525  Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Gly Gly Gly Asp Pro Val Ser Gly Pro Trp 560  Asp Pro Arg Ala Ala Pro Leu Leu Lau Arg Pro Val Gly Pro Leu Pro Trp 560  Arg Pro Arg Arg Arg His Val Ala Arg Val Ala Arg Glu Gly Gly Gly Arg Arg Glu Gln Gly Thr Val 565  Leu Leu Leu Leu Trp Ser Gly Ala Arg Val Ala Arg Pro Val Gly Pro Leu Pro Trp 560  Asp Pro Arg Ala Ala Pro Leu Leu Ala Leu Arg Arg Leu Arg Glu Gln Gly Thr Val 565  Arg Pro Leu Leu Leu Leu Leu Leu Leu Ala Leu Arg Arg Leu Leu Leu His Ala Arg Pro Arg Pro Arg Pro Ser Gly Ala Ala Arg Pro Ser Arg Leu Cys Arg Arg Pro Ser Arg Pro Leu	Leu	Pro	Asn	Ala	Thr 320	Ala	Arg	Glu	Ser	Asp 325	Gly	Trp	Tyr	Val	Leu 330
Thr Ser Trp Asn Val Ser Met Asp Thr Gln Ala Gln Gln Leu Ile 375  Leu His Phe Ser Ser Arg Met His Ala Thr Phe Ser Ala Ala Trp 380  Ser Leu Pro Gly Leu Gly Gln Asp Thr Leu Val Pro Pro Val Tyr 405  Thr Val Ser Gln Ala Arg Gly Ser Ser Pro Val Ser Leu Asp Leu 415  Ser Asp Val Gln Phe Ala Trp Lys His Leu Leu Cys Pro Asp Val Ser Tyr Arg His Leu Gly Leu Leu His Afo	Glu	Lys	Val	Asp		His	Pro	Gln	Leu	Cys 340	Phe	Lys	Phe	Ser	Phe 345
See   Heu   His   Phe   See   See   Arg   Met   His   Ala   Thr   385   Ree   See   Ala   Ala   Thr   385   Ree   See   Ala   Ala   Thr   385   Ree   See   Ala   Ala   Thr   385   Ree   Ala   Ala	Gly	Asn	Ser	Ser		Val	Glu	Cys	Pro	His 355	Gln	Thr	Gly	Ser	Leu 360
Ser Leu         Pro         Gly         Leu and same	Thr	Ser	Trp	Asn		Ser	Met	Asp	Thr	Gln 370	Ala	Gln	Gln	Leu	Ile 375
Thr Val Ser Gln Ala Arg Gly Ser Ser Pro Val Ser Leu Asp Leu Ado Ser Asp Val Gln Phe Ado And Arg Cly Leu Cys Cys Ado Ser Cys Val Leu Cys Pro Asp Ado Ser Tyr Arg His Leu Gly Val Val Leu Ala Leu Ado Ado And Arg	Leu	His	Phe	Ser	Ser 380	Arg	Met	His	Ala	Thr 385	Phe	Ser	Ala	Ala	Trp 390
11e   11e   Pro   Pro   Leu   Arg   Pro   Gly   Cys   Cys   Val   Leu   Val   Trp   Arg   Arg	Ser	Leu	Pro	Gly		Gly	Gln	Asp	Thr	Leu 400	Val	Pro	Pro	Val	Tyr 405
Ser         Asp         Val         Gln         Phe A40         Ala         Trp         Lys         His         Leu         Leu         Cys         Pro         Asp         Val 450           Ser         Tyr         Arg         His         Leu 400         Leu         Leu         Ile         Leu 460         Ala         Leu         Leu         Ala         Leu         Ala         Leu 465           Leu         Thr         Leu         Gly         Val         Val         Leu         Ala         Leu         Leu         Arg         Pro 480           Gln         Ser         Gly         Pro         Gly         Pro         Ala         Arg         Pro         Val         Leu         Leu         Leu         Leu         His         Ala         Arg         Pro         Val         Leu         Leu         His         Ala         Arg         Arg         Leu         Ala         Arg         His         Val         Ala         Arg         Ala         Ala         Ala         Arg         Fro         Ala         Arg         Pro         Ala         Arg         Pro         Trp         Arg         Fro         Trp         Arg         Val         Ala	Thr	Val	Ser	Gln		Arg	Gly	Ser	Ser	Pro 415	Val	Ser	Leu	Asp	Leu 420
Ser         Tyr         Arg         His         Leu 455         Leu Leu Leu Leu 465         Leu Leu 465         Leu 465         Leu 465         Leu 466         Ala Leu 465         Arg         Leu Arg	Ile	Ile	Pro	Phe		Arg	Pro	Gly	Cys	Cys 430	Val	Leu	Val	Trp	Arg 435
Leu Thr Leu Leu Gly 470 Val Val Leu Ala Leu Thr Cys Arg Arg Pro 480  Gln Ser Gly Pro Gly 485 Pro Ala Arg Pro Val Leu Leu Leu His Ala 495  Ala Asp Ser Glu Ala Gln Arg Arg Leu Val Gly Ala Leu Ala Glu 510  Leu Leu Arg Ala Ala Leu Gly Gly Gly Gly Arg Asp Val Ile Val Asp 525  Leu Trp Glu Gly Arg 530 Pro Arg Arg Val Ala Arg Gly Pro Leu Pro 540  Leu Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Gly Gly Gly Gly Gly Gly Gly Fro Leu Pro 540  Leu Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Gly Gly Gly Gly Gly Gly Fro Leu Pro 540  Asp Pro Arg Ala Ala Pro Leu Leu Arg Val Ala Arg Gly	Ser	Asp	Val	Gln		Ala	Trp	Lys	His	Leu 445	Leu	Cys	Pro	Asp	Val 450
Gln Ser Gly Pro Gly Ala Arg Pro Val Leu Leu His Ala Arg Sol	Ser	Tyr	Arg	His		Gly	Leu	Leu	Ile	Leu 460	Ala	Leu	Leu	Ala	Leu 465
Ala Asp Ser Glu Ala Gln Arg Arg Leu Val Gly Ala Leu Ala Glu 500  Leu Leu Arg Ala Ala Leu Gly Gly Gly Arg Asp Val Ile Val Asp 525  Leu Trp Glu Gly Arg His Val Ala Arg Val Gly Fro Leu Pro Trp 540  Leu Trp Ala Ala Arg Thr Arg Val Ala Arg 550  Leu Leu Leu Leu Trp Ser 560  Asp Pro Arg Ala Ala Pro Leu Leu Ala Arg Pro Leu His Ala Ala Pro 585  Arg Pro Leu Leu Leu Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600	Leu	Thr	Leu	Leu		Val	Val	Leu	Ala	Leu 475	Thr	Cys	Arg	Arg	Pro 480
Leu Leu Arg Ala Ala Leu Gly Gly Gly Arg Asp Val Ile Val Asp 525  Leu Trp Glu Gly Arg His Val Ala Arg Val Gly Pro Leu Pro 540  Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Glu Glu Gln Gly Thr Val 555  Leu Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 570  Asp Pro Arg Ala Ala Pro Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600	Gln	Ser	Gly	Pro		Pro	Ala	Arg	Pro	Val 490	Leu	Leu	Leu	His	Ala 495
Leu Leu Arg Ala Ala Leu Gly Gly Gly Arg Asp Val Ile Val Asp 525  Leu Trp Glu Gly Arg His Val Ala Arg Val Gly Pro Leu Pro Trp 540  Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Gly Gly Gly Gly Gly Pro Leu Pro 540  Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 555  Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 570  Asp Pro Arg Ala Ala Pro Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600	Ala	Asp	Ser	Glu		_Gln	Arg	Arg	_Leu	Val	Gly	Ala	Leu	Ala	Glu 510
Leu Trp Glu Gly Arg His Val Ala Arg Val Gly Pro Leu Pro Trp 540  Leu Trp Ala Ala Arg Thr Arg Val Ala Arg Glu Gln Gly Thr Val 555  Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 570  Asp Pro Arg Ala Ala Pro Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600							_					**- 7	T] ^	170 l	7 an
Leu Trp Ala Ala Arg Thr Arg Val Ala Arg 550 Glu Gln Gly Thr Val 555  Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 560  Asp Pro Arg Ala Ala Pro Leu Leu Leu Ala Leu Leu His Ala Ala Pro 585  Arg Pro Leu Leu Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600					515					520					525
Leu Leu Leu Trp Ser Gly Ala Asp Leu Arg Pro Val Ser Gly Pro 560  Asp Pro Arg Ala Ala Pro Leu Leu Ala Leu Leu His Ala Ala Pro 585  Arg Pro Leu Leu Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 600					530					535					540
Asp Pro Arg Ala Ala Pro Leu Leu Ala Leu Leu His Ala Ala Pro 585  Arg Pro Leu Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 590	Leu	Trp	Ala	Ala			Arg	Val	Ala	Arg 550	Glu	Gln	Gly	Thr	Val 555
Arg Pro Leu Leu Leu Ala Tyr Phe Ser Arg Leu Cys Ala Lys 590 595 600	Leu	Leu	Leu	Trp			Ala	Asp	Leu	Arg 565	Pro	Val	Ser	Gly	Pro 570
590 595	Asp	Pro	Arg	, Ala			Lev	. Leu	Ala	Leu 580	Leu	His	Ala	a Ala	Pro 585
Gly Asp Ile Pro Pro Pro Leu Arg Ala Leu Pro Arg Tyr Arg Leu	Arg	Pro	Leu	ı Leu			ı Ala	Tyr	Phe	Ser 595	Arg	Leu	Cys	s Ala	600
	 Gly	Asp	Ile	Pro	Pro	Pro	Lev	a Arg	Ala	Leu	Pro	Arg	Туг	Arg	Leu

605 . 610 615
Leu Arg Asp Leu Pro Arg Leu Leu Arg Ala Leu Asp Ala Arg Pro 620 625 630
Phe Ala Glu Ala Thr Ser Trp Gly Arg Leu Gly Ala Arg Gln Arg 635 640 645
Arg Gln Ser Arg Leu Glu Leu Cys Ser Arg Leu Glu Arg Glu Ala 650 655 660
Ala Arg Leu Ala Asp Leu Gly 665
<210> 17 <211> 2319 <212> DNA <213> Homo Sapien
<400> 17 gccaggccct atctccctgc caggaggccg gagtggggga ggtcagacgg 50
ggcggttgga gggggaggga tgccacgcgc ttctgcctca ggtgttcctg 100
cgttgtttgt cagtggagag cagggagtgg ggccagccag cagaaacagt 150
gggctgtaca acatcacctt caaatatgac aattgtacca cctacttgaa 200
tccagtgggg aagcatgtga ttgctgacgc ccagaatatc accatcagcc 250
agtatgettg ccatgaccaa gtggcagtca ccattetttg gteeccaggg 300
gccctcggca tcgaattcct gaaaggattt cgggtaatac tggaggagct 350
gaagtcggag ggaagacagt gccaacaact gattctaaag gatccgaagc 400
agctcaacag tagcttcaaa agaactggaa tggaatctca acctttcctg 450
aatatgaaat ttgaaacgga ttatttcgta aaggttgtcc cttttccttc 500
cattaaaaac gaaagcaatt accacccttt cttctttaga acccgagcct 550
gtgacctgtt gttacagccg gacaatctag cttgtaaacc cttctggaag 600
cctcggaacc tgaacatcag ccagcatggc tcggacatgc aggtgtcctt 650

cgaccacgca ccgcatggct cggacatgca ggtgtccttc gaccacgcac 700

cgcacaactt cggcttccgt ttcttctatc ttcactacaa gctcaagcac 750

gaaggacctt tcaagcgaaa gacctgtaag caggagcaaa ctacagagat 800

gaccagetge etectteaaa atgtttetee aggggattat ataattgage 850

tggtggatga cactaacaca acaagaaaag tgatgcatta tgccttaaag 900

ccagtgcact ccccgtgggc cgggcccatc agagccgtgg ccatcacagt 950

gccactggta	gtcatatcgg	cattcgcgac	gctcttcact	gtgatgtgcc	1000
gcaagaagca	acaagaaaat	atatattcac	atttagatga	agagagctct	1050
gagtcttcca	catacactgc	agcactccca	agagagaggc	tccggccgcg	1100
gccgaaggtc	tttctctgct	attccagtaa	agatggccag	aatcacatga	1150
atgtcgtcca	gtgtttcgcc	tacttcctcc	aggacttctg	tggctgtgag	1200
gtggctctgg	acctgtggga	agacttcagc	ctctgtagag	aagggcagag	1250
agaatgggtc	atccagaaga	tccacgagtc	ccagttcatc	attgtggttt	1300
gttccaaagg	tatgaagtac	tttgtggaca	agaagaacta	caaacacaaa	1350
ggaggtggcc	gaggctcggg	gaaaggagag	ctcttcctgg	tggcggtgtc	1400
agccattgcc	gaaaagctcc	gccaggccaa	gcagagttcg	tccgcggcgc	1450
tcagcaagtt	tatcgccgtc	tactttgatt	attcctgcga	gggagacgtc	1500
cccggtatcc	tagacctgag	taccaagtac	agactcatgg	acaatcttcc	1550
tcagctctgt	tcccacctgc	actcccgaga	ccacggcctc	caggagccgg	1600
ggcagcacac	gcgacagggc	agcagaagga	actacttccg	gagcaagtca	1650
ggccggtccc	tatacgtcgc	catttgcaac	atgcaccagt	ttattgacga	1700
ggagcccgac	tggttcgaaa	agcagttcgt	tcccttccat	cctcctccac	1750
tgcgctaccg	ggagccagtc	ttggagaaat	ttgattcggg	cttggtttta	1800
aatgatgtca	tgtgcaaacc	agggcctgag	agtgacttct	gcctaaaggt	1850
agaggcggct	gttcttgggg	caaccggacc	agccgactcc	cagcacgaga	1900
gtcagcatgg	gggcctggac	caagacgggg	aggcccggcc	tgcccttgac	1950
ggtagcgccg	ccctgcaacc	cctgctgcac	acggtgaaag	ccggcagccc	2000
ctcggacatg	ccgcgggact	caggcatcta	tgactcgtct	gtgccctcat	2050
ccgagctgtc	tctgccactg	atggaaggac	tctcgacgga	ccagacagaa	2100
acgtcttccc	tgacggagag	cgtgtcctcc	tcttcaggcc	tgggtgagga	2150
ggaacctcct	gcccttcctt	ccaagctcct	ctcttctggg	tcatgcaaag	2200
cagatcttgg	ttgccgcagc	tacactgatg	aactccacgc	ggtcgcccct	2250
ttgtaacaaa	acgaaagagt	ctaagcattg	ccactttaaa	aaaaaaaaaa	2300
aaaaaaaaa	aaaaaaaaa :	2319			

<sup>&</sup>lt;210> 18 <211> 728

.<212>	PRT.	
<213>	Homo	Sapien

•	<400: Met 1		Arg	Ala	Ser 5	Ala	Ser	Gly	Val	Pro 10	Ala	Leu	Phe	Val	Ser 15
	Gly	Glu	Gln	Gly	Val 20	Gly	Pro	Ala	Ser	Arg 25	Asn	Ser	Gly	Leu	Tyr 30
	Asn	Ile	Thr	Phe	Lys 35	Tyr	Asp	Asn	Cys	Thr 40	Thr	Tyr	Leu	Asn	Pro 45
	Val	Gly	Lys	His	Val 50	Ile	Ala	Asp	Ala	Gln 55	Asn	Ile	Thr	Ile	Ser 60
	Gln	Tyr	Ala	Cys	His 65	Asp	Gln	Val	Ala	Val 70	Thr	Ile	Leu	Trp	Ser 75
	Pro	Gly	Ala	Leu	Gly 80	Ile	Glu	Phe	Leu	Lys 85	Gly	Phe	Arg	Val	Ile 90
	Leu	Glu	Glu	Leu	Lys 95	Ser	Glu	Gly	Arg	Gln 100	Cys	Gln	Gln	Leu	Ile 105
	Leu	Lys	Asp	Pro	Lys 110	Gln	Leu	Asn	Ser	Ser 115	Phe	Lys	Arg	Thr	Gly 120
	Met	Glu	Ser	Gln	Pro 125	Phe	Leu	Asn	Met	Lys 130	Phe	Glu	Thr	Asp	Tyr 135
	Phe	Val	Lys	Val	Val 140	Pro	Phe	Pro	Ser	Ile 145	Lys	Asn	Glu	Ser	Asn 150
	Tyr	His	Pro	Phe	Phe 155	Phe	Arg	Thr	Arg	Ala 160	Cys	Asp	Leu	Leu	Leu 165
	Gln	Pro	Asp	Asn	Leu 170	Ala	Cys	Lys	Pro	Phe 175	Trp	Lys	Pro	Arg	Asn 180
	Leu	Asn	Ile	Ser	Gln 185	His	Gly	Ser	Asp	Met 190	Gln	Val	Ser	Phe	Asp 195
	His	Ala	Pro	His	Gly 200	Ser	Asp	Met	Gln	Val 205	Ser	Phe	Asp	His	Ala 210
	Pro	His	Asn	Phe	Gly 215	Phe	Arg	Phe	Phe	Tyr 220	Leu	His	Tyr	Lys	Leu 225
	Lys	His	Glu	Gly	Pro 230	Phe	Lys	Arg	Lys	Thr 235	Cys	Lys	Gln	Glu	Gln 240
	Thr	Thr	Glu	Met	Thr 245	Ser	Cys	Leu	Leu	Gln 250	Asn	Val	Ser	Pro	Gly 255
	Asp	Tyr	Ile	Ile	Glu 260	Leu	Val	Asp	Asp	Thr 265	Asn	Thr	Thr	Arg	Lys 270

Val	Met	His	Tyr	Ala 275	Leu	Lys	Pro	Val	His 280	Ser	Pro	Trp	Ala	Gly 285
Pro	Ile	Arg	Ala	Val 290	Ala	Ile	Thr	Val	Pro 295	Leu	Val	Val	Ile	Ser 300
Ala	Phe	Ala	Thr	Leu 305	Phe	Thr	Val	Met	Cys 310	Arg	Lys	Lys	Gln	Gln 315
Glu	Asn	Ile	Tyr	Ser 320	His	Leu	Asp	Glu	Glu 325	Ser	Ser	Glu	Ser	Ser 330
Thr	Tyr	Thr	Ala	Ala 335	Leu	Pro	Arg	Glu	Arg 340	Leu	Arg	Pro	Arg	Pro 345
Lys	Val	Phe	Leu	Cys 350	Tyr	Ser	Ser	Lys	Asp 355	Gly	Gln	Asn	His	Met 360
Asn	Val	Val	Gln	Cys 365	Phe	Ala	Tyr	Phe	Leu 370	Gln	Asp	Phe	Cys	Gly 375
Cys	Glu	Val	Ala	Leu 380	Asp	Leu	Trp	Glu	Asp 385	Phe	Ser	Leu	Cys	Arg 390
Glu	Gly	Gln	Arg	Glu 395	Trp	Val	Ile	Gln	Lys 400	Ile	His	Glu	Ser	Gln 405
Phe	Ile	Ile	Val	Val 410	Cys	Ser	Lys	Gly	Met 415	Lys	Tyr	Phe	Val	Asp 420
Lys	Lys	Asn	Tyr	Lys 425	His	Lys	Gly	Gly	Gly 430	Arg	Gly	Ser	Gly	Lys 435
Gly	Glu	Leu	Phe	Leu 440	Val	Ala	Val	Ser	Ala 445	Ile	Ala	Glu	Lys	Leu 450
Arg	Gln	Ala	Lys	Gln 455	Ser	Ser	Ser	Ala	Ala 460	Leu	Ser	Lys	Phe	Ile 465
Ala	Val	Tyr			_	Ser	Cys	Glu		_	Val	Pro	_	
Leu	Asp	Leu	Ser	Thr 485	Lys	Tyr	Arg	Leu	Met 490	Asp	Asn	Leu	Pro	Gln 495
Leu	Cys	Ser	His	Leu 500	His	Ser	Arg	Asp	His 505	Gly	Leu	Gln	Glu	Pro 510
Gly	Gln	His	Thr	Arg 515	Gln	Gly	Ser	Arg	Arg 520	Asn	Tyr	Phe	Arg	Ser 525
Lys	Ser	Gly	Arg	Ser 530	Leu	Tyr	Val	Ala	Ile 535	Cys	Asn	Met	His	Gln 540
Phe	Ile	Asp	Glu	Glu 545	Pro	Asp	Trp	Phe	Glu 550	Lys	Gln	Phe	Val	Pro 555
Phe	His	Pro	Pro	Pro	Leu	Arg	Tyr	Arg	Glu	Pro	Val	Leu	Glu	Lys

		·		560					565					570
Phe	Asp	Ser	Gly	Leu 575	Val	Leu	Asn	Asp	Val 580	Met	Cys	Lys	Pro	Gly 585
Pro	Glu	Ser	Asp	Phe 590	Cys	Leu	Lys	Val	Glu 595	Ala	Ala	Val	Leu	Gly 600
Ala	Thr	Gly	Pro	Ala 605	Asp	Ser	Gln	His	Glu 610	Ser	Gln	His	Gly	Gly 615
Leu	Asp	Gln	Asp	Gly 620	Glu	Ala	Arg	Pro	Ala 625	Leu	Asp	Gly	Ser	Ala 630
Ala	Leu	Gln	Pro	Leu 635	Leu	His	Thr	Val	Lys 640	Ala	Gly	Ser	Pro	Ser 645
Asp	Met	Pro	Arg	Asp 650	Ser	Gly	Ile	Tyr	Asp 655	Ser	Ser	Val	Pro	Ser 660
Ser	Glu	Leu	Ser	Leu 665	Pro	Leu	Met	Glu	Gly 670	Leu	Ser	Thr	Asp	Gln 675
Thr	Glu	Thr	Ser	Ser 680	Leu	Thr	Glu	Ser	Val 685	Ser	Ser	Ser	Ser	Gly 690
Leu	Gly	Glu	Glu	Glu 695	Pro	Pro	Ala	Leu	Pro 700	Ser	Lys	Leu	Leu	Ser 705
Ser	Gly	Ser	Cys	Lys 710	Ala	Asp	Leu	Gly	Cys 715	Arg	Ser	Tyr	Thr	Asp 720
Glu	Leu	His	Ala	Val 725	Ala	Pro	Leu							
	<210> 19 <211> 24													
<212>	> DNA		,; , ]	Com	onac									
		.111	Jai	sequ	ience	•								
<220> <223>		thet	ic C	ligo	nucl	eoti	de F	robe	<b>!</b>					
<400> 19 atccacagaa gctggccttc gccg 24														
<210> 20 <211> 24 <212> DNA <213> Artificial Sequence														
<220> <223> Synthetic Oligonucleotide Probe														
<400> ggga		ga t	gaac	tcgg	t gt	gg 2	4							

<210> 21

```
<211> 40 .
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 21
 tatccacaga agctggcctt cgccgagtgc ctgtgcagag 40
<210> 22
<211> 27
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 22
 gttgcattct tggcaatggt catggga 27
<210> 23
<211> 24
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 23
 ggtccatgtg ggagcctgtc tgta 24
<210> 24
<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 24
cagcagetee teagaggtgt cetgeeettt getggggeag eaget 45
<210> 25
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 25
gctcagtgcc ttccaccaca cgc 23
<210> 26
<211> 21
<212> DNA
<213> Artificial Sequence
```

```
<220>
<223> Synthetic Oligonucleotide Probe
<400> 26
 ctgcgtcctt ctccggctcg g 21
<210> 27
<211> 45
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 27
 cgttccgtct acaccgaggc ctacgtcacc atccccgtgg gctgc 45
<210> 28
<211> 25
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 28
 actccatatt ttcctacttg tggca 25
<210> 29
<211> 19
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 29
 cccaaagtga cctaagaac 19
<210> 30
<211> 27
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 30
tcactgaatt tcttcaaaac cattgca 27
<210> 31
<211> 50
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
```

```
<400> 31
 tgtggcagcg actgcatccg acataaagga acagttgtgc tctgcccaca 50
<210> 32
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 32
 ccgacttctt gcagggccgg 20
<210> 33
<211> 20
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 33
 gcagcacgca gctgagcgag 20
<210> 34
<211> 33
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 34
 agcgagtggc tacaggatgg ggtgtccggg ccc 33
<210> 35
<211> 25
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 35
 cgttgtttgt cagtggagag caggg 25
<210> 36
<211> 23
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 36
caggaacacc tgaggcagaa gcg 23
```

```
<210> 37
<211> 40
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 37
 ctatctccct gccaggaggc cggagtgggg gaggtcagac 40
<210> 38
<211> 21
<212> DNA
<213> Artificial Sequence
<223> Synthetic Oligonucleotide Probe
<400> 38
ctgtacctcg agggtgcaga g 21
<210> 39
<211> 58
<212> DNA
<213> Artificial Sequence
<220>
<223> Synthetic Oligonucleotide Probe
<400> 39
cccaagettg ggtcaatgat gatgatgatg atgatgatgc cacaggggca 50
 tgtagtcc 58
```